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CYTOKINE DESIGNATED LERK-6

Abstract:

The invention is directed to LERK-6 as a purified and isolated protein, the DNA encoding the LERK-6, host cells transfected with cDNAs encoding LERK-6.

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TITLE
CYTOKINE DESIGNATED LERK-6
RELATED U.S. APPLICATION DATA

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This is a continuation-in-part of U.S. Ser. No. 08/538,709, filed October 3, 1995, currently pending, which is a continuation-in-part of U.S. Ser. No. 08/318,393, filed October 5, 1994.

FIELD OF THE INVENTION

10

The present invention relates to cytokine polypeptides designated as LERK-6 that bind to the hek or elk receptor, the nucleic acids encoding such polypeptides, processes for production of recombinant LERK-6 polypeptides, and pharmaceutical compositions containing such polypeptides.

15

BACKGROUND OF THE INVENTION

Proteins known as the receptor tyrosine kinases have an intrinsic kinase activity that is activated upon ligand binding. This class of proteins is characterized by conserved structural motifs within the catalytic domains (Hanks et al., *Science*, 242:42, 1988) and can be subdivided into families based on structural features of the regions N-terminal to the catalytic domain.

Boyd et al. (*J. Biol. Chem.*, 267:3262, 1992) purified a cell-surface glycoprotein exhibiting tyrosine kinase activity. The amino acid sequence identified this protein as a member of the *eph/elk* family, and the protein was thus designated hek (human *eph/elk*-like kinase). A monoclonal antibody immunoreactive with hek was used to study hek expression on a number of human cell types (Boyd et al., *supra*). Hek antigen was detected on the human pre-B cell leukemia cell line LK63 (the cell line employed as the immunogen against which the antibody was raised) and the human T-cell leukemia cell line, JM. The Raji B lymphoma cell line showed weak hek antigen expression, and the remaining cell lines tested (both normal and tumor cell lines, among which were hemopoietic cell lines that included pre-B and T-cell lines) were consistently negative. Of the normal and tumor tissue biopsy specimens that were also tested for hek antigen expression, none of the normal tissues was positive and only a very low proportion of hemopoietic tumors was positive.

35

Expression of hek transcripts in the above-described LK63 and JM cell lines, as well as the human T-cell leukemia cell line HSB-2, has been demonstrated by northern blot analysis (Wicks et al., *Proc. Natl. Acad. Sci. USA*, 89:1611, 1992). Nucleotide and

amino acid sequences for an isolated hek cDNA clone are presented in Wicks et al., *supra*.

5 The cell surface protein designated elk is another member of the eph-related tyrosine kinase receptor family of proteins. A partial clone of elk was first discovered in a rat brain cDNA expression library that was screened for proteins expressing tyrosine kinase activity (Letwin et al., *Oncogene* 3:621, 1988). Later, a composite sequence spanning the entire elk coding region was derived from partial clones isolated from a rat brain cDNA library and a rat cerebellar brain library using the partial clone as a probe
10 (Lhotak et al., *Mol. Cell. Biol.* 11:2496, 1991).

The hek and elk proteins are closely related to a number of other receptor tyrosine kinases, including the hek homologs mek4 and cek4 (Sajjadi et al. *New Biol.* 3:769, 1991); eek (Chan et al. *Oncogene* 6:1057, 1991); erk (Chan et al. *supra*), eck (Lindberg
15 et al. *Mol. Cell. Biol.* 10:6316, 1990); cek5 (Pasquale, E.B. *Cell Regulation* 2:523, 1991); and eph (Hirai et al. *Science* 238:1717, 1987). The proteins of this subfamily are related not only in their cytoplasmic domains, but also in their extracellular domains, which are 41 to 68% identical. Interestingly, the tissue distributions of these various receptors are diverse. For example, expression of elk mRNA has been reported to be limited to testis
20 and brain (Lhotak et al., *supra*), whereas eck is found not only in these same two tissues but in lung, intestine, kidney, spleen, ovary, and skin as well. In addition, most eph-related receptors are primarily expressed in the brain. Due to the homology of the receptors in the eph family, a given ligand for one specific receptor may also bind other receptors.

25 Those ligands that have been identified for the receptor tyrosine kinases are a diverse group of proteins that affect the growth, differentiation, and survival of cells expressing the receptors. Ligands for hek and elk have been isolated, as discussed in more detail below.

30 Identification of additional ligands for hek and elk that may exist would prove useful in investigating the nature of cellular processes regulated by signaling through these receptors. If enhancement or inhibition of a particular biological signal mediated through these receptors is desired, it is advantageous to identify each of the proteins that
35 may play a role in transduction of such signals. Further, it is known that certain proteins can bind to receptors without initiating signal transduction, including interleukin-1 receptor antagonist protein (Eisenberg et al., *Nature* 343:341, 1990; Hannum et al.,

Nature 343:336, 1990; and Carter et al., *Nature* 344:633, 1990). Identification of additional proteins that bind hek or elk is also desirable in order to determine whether such proteins function as antagonists.

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SUMMARY OF THE INVENTION

The present invention pertains to mammalian LERK-6 as an isolated or homogeneous protein. In addition, the invention is directed to isolated DNAs encoding mammalian LERK-6 and to expression vectors comprising a cDNA encoding mammalian LERK-6. Within the scope of this invention are host cells that have been transfected or
10 transformed with expression vectors that comprise a cDNA encoding LERK-6, and processes for producing LERK-6 by culturing such host cells under conditions conducive to expression of LERK-6.

In addition, LERK-6 can be bound to a solid phase matrix and used to affinity-
15 purify or separate cells that express hek or elk on their cell surface. The invention encompasses separating cells having the hek or elk receptor on the surface thereof from a mixture of cells in solution, comprising contacting the cells in the mixture with a contacting surface having a LERK-6 molecule thereon, and separating the contacting
20 surface and the solution.

20

DETAILED DESCRIPTION OF THE INVENTION

A cDNA encoding murine LERK-6 has been isolated and is disclosed in SEQ ID NO:1. An exon from the human LERK-6 has been isolated and is disclosed in SEQ ID
25 NO:7. DNA encoding a full length human LERK-6 has been isolated and is disclosed in SEQ ID NO:9. This discovery of a cDNA encoding LERK-6 enables construction of expression vectors comprising cDNAs encoding LERK-6; host cells transfected or transformed with the expression vectors; biologically active LERK-6 as homogeneous proteins; and antibodies immunoreactive with LERK-6.

30

LERK-6 may be useful in the enhancement, stimulation, proliferation or growth of cells expressing the hek or elk receptor. Since the hek or elk receptor is found in the tissue of the brain and testis, treatment of a variety of conditions associated with tissue damage thereof is possible. Moreover, the ligand and receptor complex may be involved
35 in neural growth, development and/or maintenance. While not limited to such, particular uses of the LERK-6 are described infra.

As used herein, the term "LERK-6" refers to a genus of polypeptides that bind and complex independently with hek or elk receptor found on T-cells and brain cells. The term "LERK-6" encompasses polypeptides having the amino acid sequence 1-184 of SEQ ID NO:2, proteins that are encoded by nucleic acids that contain the nucleic acid sequence of SEQ ID NO:7, and polypeptides having the amino acid sequence 1 -184 of SEQ ID NO:10. In addition, LERK-6 encompasses polypeptides that have a high degree of similarity or a high degree of identity with the amino acid sequence 1-184 of SEQ ID NO:2, the amino acid sequence of SEQ ID NO:8, and amino acids 1-184 of SEQ ID NO:10, and which polypeptides are biologically active and bind the hek or elk receptor. In addition, the term "murine LERK-6" refers to biologically active gene products of the DNA of SEQ ID NO:1 and the term "human LERK-6" refers to biologically active gene products of the DNA of SEQ ID NO:9.. Further encompassed by the term "LERK-6" are the GPI-linked proteins (which include an extracellular region and a C-terminal hydrophobic region), and soluble or truncated proteins that comprise primarily the receptor-binding portion of the protein, retain biological activity and are capable of being secreted. Specific examples of such soluble proteins are those comprising the sequence of amino acids 1 (Ala) - 145 (Asn) of SEQ ID NO:2 and those comprising the sequence of amino acids 1-145 of SEQ ID NO:10. .

The term "hek/elk" means either hek or elk or both hek and elk. For example, the term "anti-hek/elk antibodies" refers to antibodies against either hek or elk. The term "hek/elk-expressing cells" refers to cells that express either the hek receptor or the elk receptor, or cells that express both the hek and elk receptors.

The term "biologically active" as it refers to LERK-6, means that the LERK-6 is capable of binding to hek/elk.

"Isolated" means that LERK-6 is free of association with other proteins or polypeptides, for example, as a purification product of recombinant host cell culture or as a purified extract.

A "LERK-6 variant" as referred to herein, means a polypeptide substantially homologous to native LERK-6, but which has an amino acid sequence different from that of native LERK-6 (human, murine or other mammalian species) because of one or more deletions, insertions or substitutions. The variant amino acid sequence preferably is at least 80% identical to a native LERK-6 amino acid sequence, most preferably at least 90% identical. The percent identity may be determined, for example, by comparing

sequence information using the GAP computer program, version 6.0 described by Devereux et al. (*Nucl. Acids Res.* 12:387, 1984) and available from the University of Wisconsin Genetics Computer Group (UWGCG). The preferred default parameters for the GAP program include: (1) a unary comparison matrix (containing a value of 1 for identities and 0 for non-identities) for nucleotides, and the weighted comparison matrix of Gribkov and Burgess, *Nucl. Acids Res.* 14:6745, 1986, as described by Schwartz and Dayhoff, eds., *Atlas of Protein Sequence and Structure*, National Biomedical Research Foundation, pp. 353-358, 1979; (2) a penalty of 3.0 for each gap and an additional 0.10 penalty for each symbol in each gap; and (3) no penalty for end gaps. Variants may comprise conservatively substituted sequences, meaning that a given amino acid residue is replaced by a residue having similar physiochemical characteristics. Examples of conservative substitutions include substitution of one aliphatic residue for another, such as Ile, Val, Leu, or Ala for one another, or substitutions of one polar residue for another, such as between Lys and Arg; Glu and Asp; or Gln and Asn. Other such conservative substitutions, for example, substitutions of entire regions having similar hydrophobicity characteristics, are well known. Naturally occurring LERK-6 variants or alleles are also encompassed by the invention. Examples of such variants are proteins that result from alternate mRNA splicing events or from proteolytic cleavage of the LERK-6 protein, wherein the LERK-6 binding property is retained. Alternate splicing of mRNA may yield a truncated but biologically active LERK-6 protein, such as a naturally occurring soluble form of the protein, for example. Variations attributable to proteolysis include, for example, differences in the N- or C-termini upon expression in different types of host cells, due to proteolytic removal of one or more terminal amino acids from the LERK-6 protein (generally from 1-5 terminal amino acids).

LERK-6 is predicted to be anchored to the cell surface via glycosylphosphatidylinositol (GPI) linkage. GPI membrane anchors, including the chemical structure and processing thereof, are described in Ferguson, M. and Williams, A., *Ann. Rev. Biochem.*, 57:285, 1988 (hereby incorporated by reference). When initially expressed, certain proteins comprise a C-terminal hydrophobic domain that contains signals for GPI anchoring. A cleavage site is located upstream, often about 10-12 amino acids upstream of the N-terminus of the hydrophobic domain. Post translational processing includes cleavage of the protein at this cleavage site. A GPI anchor attaches to the newly exposed C-terminal amino acid of the processed, mature protein. Thus, when LERK-6 proteins are expressed in cells that recognize GPI anchoring signals in the hydrophobic domain, the full length amino acid sequence of SEQ ID NO:2 and SEQ ID NO:10 represents precursor forms of the proteins.

Based on consensus sequences derived from other GPI-anchored proteins, the hydrophobic region in LERK-6 of the present invention likely is between and includes, amino acids 170 (Leu) and 184 (Ser) of SEQ ID NO:2 and SEQ ID NO:10. Between the hydrophobic domain and the receptor-binding domain, is a spacer region of about 22 to 25 amino acids in length. The spacer region extends from approximately amino acid 145 (Asn), 146 (Glu) or 147 (Thr) to about amino acid 169 (His) of SEQ ID NO:2 and SEQ ID NO:10

Example 3 describes the construction of a novel hek:Fc fusion protein that may be utilized in the screening for LERK-6. Other antibody Fc regions may be substituted for the human IgG1 Fc region described in the Example. Other suitable Fc regions are those that can bind with high affinity to protein A or protein G, and include the Fc region of human IgG1 or fragments of the human or murine IgG1 Fc region, e.g., fragments comprising at least the hinge region so that interchain disulfide bonds will form. The hek:Fc fusion protein offers the advantage of being easily purified. In addition, disulfide bonds form between the Fc regions of two separate fusion protein chains, creating dimers.

As described supra, an aspect of the invention is soluble LERK-6 polypeptides. Soluble LERK-6 polypeptides comprise all or part of the extracellular domain of a native LERK-6 but lack the GPI signal that would cause retention of the polypeptide on a cell membrane. Soluble LERK-6 polypeptides advantageously comprise the native (or a heterologous) signal peptide when initially synthesized to promote secretion, but the signal peptide is cleaved upon secretion of LERK-6 from the cell. Soluble LERK-6 polypeptides encompassed by the invention retain the ability to bind the hek or elk receptor. Indeed, soluble LERK-6 may also include part of the GPI signal or part of the cytoplasmic domain or other sequences, provided that the soluble LERK-6 protein can be secreted.

Soluble LERK-6 may be identified (and distinguished from its non-soluble membrane-bound counterparts) by separating intact cells which express the desired protein from the culture medium, e.g., by centrifugation, and assaying the medium (supernatant) for the presence of the desired protein. The presence of LERK-6 in the medium indicates that the protein was secreted from the cells and thus is a soluble form of the desired protein.

Soluble forms of LERK-6 possess many advantages over the native bound LERK-6 protein. Purification of the proteins from recombinant host cells is feasible, since the soluble proteins are secreted from the cells. Further, soluble proteins are generally more suitable for intravenous administration.

5

Examples of soluble LERK-6 polypeptides include those comprising a substantial portion of the extracellular domain of a native LERK-6 protein. An example of a soluble murine LERK-6 protein comprises amino acids 1-145 of SEQ ID NO:2 and amino acids 1-145 of SEQ ID NO:10. In addition, truncated soluble LERK-6 proteins comprising less than the entire extracellular domain are included in the invention. When initially expressed within a host cell, soluble LERK-6 may additionally comprise one of the heterologous signal peptides described below that is functional within the host cells employed. Alternatively, the protein may comprise the native signal peptide. In one embodiment of the invention, soluble LERK-6 can be expressed as a fusion protein comprising (from N- to C-terminus) the yeast α -factor signal peptide, a FLAG[®] peptide described below and in U.S. Patent No. 5,011,912, and soluble LERK-6 consisting of amino acids 1-134 of SEQ ID NO:2 or SEQ ID NO:10. This recombinant fusion protein is expressed in and secreted from yeast cells. The FLAG[®] peptide facilitates purification of the protein, and subsequently may be cleaved from the soluble LERK-6 using bovine mucosal enterokinase. Isolated DNA sequences encoding soluble LERK-6 proteins are encompassed by the invention.

Truncated LERK-6, including soluble polypeptides, may be prepared by any of a number of conventional techniques. A desired DNA sequence may be chemically synthesized using techniques known per se. DNA fragments also may be produced by restriction endonuclease digestion of a full length cloned DNA sequence, and isolated by electrophoresis on agarose gels. Linkers containing restriction endonuclease cleavage site(s) may be employed to insert the desired DNA fragment into an expression vector, or the fragment may be digested at cleavage sites naturally present therein. The well known polymerase chain reaction procedure also may be employed to amplify a DNA sequence encoding a desired protein fragment. As a further alternative, known mutagenesis techniques may be employed to insert a stop codon at a desired point, e.g., immediately downstream of the codon for the last amino acid of the receptor-binding domain.

As stated above, the invention provides isolated or homogeneous LERK-6 polypeptides, both recombinant and non-recombinant. Variants and derivatives of native LERK-6 proteins that retain the desired biological activity (e.g., the ability to bind

hek/elk) may be obtained by mutations of nucleotide sequences coding for native LERK-6 polypeptides. Alterations of the native amino acid sequence may be accomplished by any of a number of conventional methods. Mutations can be introduced at particular loci by synthesizing oligonucleotides containing a mutant sequence, flanked by restriction sites enabling ligation to fragments of the native sequence. Following ligation, the resulting reconstructed sequence encodes an analog having the desired amino acid insertion, substitution, or deletion.

Alternatively, oligonucleotide-directed site-specific mutagenesis procedures can be employed to provide an altered gene wherein predetermined codons can be altered by substitution, deletion or insertion. Exemplary methods of making the alterations set forth above are disclosed by Walder et al. (*Gene* 42:133, 1986); Bauer et al. (*Gene* 37:73, 1985); Craik (*BioTechniques*, January 1985, 12-19); Smith et al. (*Genetic Engineering: Principles and Methods*, Plenum Press, 1981); Kunkel (*Proc. Natl. Acad. Sci. USA* 82:488, 1985); Kunkel et al. (*Methods in Enzymol.* 154:367, 1987); and U.S. Patent Nos. 4,518,584 and 4,737,462 all of which are incorporated by reference.

LERK-6 may be modified to create LERK-6 derivatives by forming covalent or aggregative conjugates with other chemical moieties, such as glycosyl groups, lipids, phosphate, acetyl groups and the like. Covalent derivatives of LERK-6 may be prepared by linking the chemical moieties to functional groups on LERK-6 amino acid side chains or at the N-terminus or C-terminus of a LERK-6 polypeptide or the extracellular domain thereof. Other derivatives of LERK-6 within the scope of this invention include covalent or aggregative conjugates of LERK-6 or its fragments with other proteins or polypeptides, such as by synthesis in recombinant culture as N-terminal or C-terminal fusions. For example, the conjugate may comprise a signal or leader polypeptide sequence (e.g. the α -factor leader of *Saccharomyces*) at the N-terminus of a LERK-6 polypeptide. The signal or leader peptide co-translationally or post-translationally directs transfer of the conjugate from its site of synthesis to a site inside or outside of the cell membrane or cell wall.

LERK-6 polypeptide fusions can comprise peptides added to facilitate purification and identification of LERK-6. Such peptides include, for example, poly-His or the antigenic identification peptides described in U.S. Patent No. 5,011,912 and in Hopp et al., *Bio/Technology* 6:1204, 1988.

The invention further includes LERK-6 polypeptides with or without associated native-pattern glycosylation. LERK-6 expressed in yeast or mammalian expression

systems (e.g., COS-7 cells) may be similar to or significantly different from a native LERK-6 polypeptide in molecular weight and glycosylation pattern, depending upon the choice of expression system. Expression of LERK-6 polypeptides in bacterial expression systems, such as *E. coli*, provides non-glycosylated molecules.

5

Equivalent DNA constructs that encode various additions or substitutions of amino acid residues or sequences, or deletions of terminal or internal residues or sequences not needed for biological activity or binding are encompassed by the invention. For example, N-glycosylation sites in the LERK-6 extracellular domain can be modified to preclude glycosylation, allowing expression of a reduced carbohydrate analog in mammalian and yeast expression systems. N-glycosylation sites in eukaryotic polypeptides are characterized by an amino acid triplet Asn-X-Y, wherein X is any amino acid except Pro and Y is Ser or Thr. The murine LERK-6 and human LERK-6 proteins comprise three such triplets, at amino acids 13-15, 145-147 and 159-161 of SEQ ID NO:2 and SEQ ID NO:10, respectively. Appropriate substitutions, additions or deletions to the nucleotide sequence encoding these triplets will result in prevention of attachment of carbohydrate residues at the Asn side chain. Alteration of a single nucleotide, chosen so that Asn is replaced by a different amino acid, for example, is sufficient to inactivate an N-glycosylation site. Known procedures for inactivating N-glycosylation sites in proteins include those described in U.S. Patent 5,071,972 and EP 276,846, hereby incorporated by reference.

In another example, sequences encoding Cys residues that are not essential for biological activity can be altered to cause the Cys residues to be deleted or replaced with other amino acids, preventing formation of incorrect intramolecular disulfide bridges upon renaturation. Other equivalents are prepared by modification of adjacent dibasic amino acid residues to enhance expression in yeast systems in which KEX2 protease activity is present. EP 212,914 discloses the use of site-specific mutagenesis to inactivate KEX2 protease processing sites in a protein. KEX2 protease processing sites are inactivated by deleting, adding or substituting residues to alter Arg-Arg, Arg-Lys, and Lys-Arg pairs to eliminate the occurrence of these adjacent basic residues. Lys-Lys pairings are considerably less susceptible to KEX2 cleavage, and conversion of Arg-Lys or Lys-Arg to Lys-Lys represents a conservative and preferred approach to inactivating KEX2 sites. The murine LERK-6 and human LERK-6 contain one KEX2 protease processing site at amino acids 81-82 of SEQ ID NO:2 and SEQ ID NO:10, respectively.

Nucleic acid sequences within the scope of the invention include isolated DNA and RNA sequences that hybridize to the LERK-6 nucleotide sequences disclosed herein under conditions of moderate or severe stringency, and that encode biologically active LERK-6. Conditions of moderate stringency, as defined by Sambrook et al. *Molecular Cloning: A Laboratory Manual*, 2 ed. Vol. 1, pp. 101-104, Cold Spring Harbor Laboratory Press, (1989), include use of a prewashing solution of 5 X SSC, 0.5% SDS, 1.0 mM EDTA (pH 8.0) and hybridization conditions of about 55°C, 5 X SSC, overnight. Conditions of severe stringency include higher temperatures of hybridization and washing. The skilled artisan will recognize that the temperature and wash solution salt concentration may be adjusted as necessary according to factors such as the length of the nucleic acid molecule.

Due to the known degeneracy of the genetic code wherein more than one codon can encode the same amino acid, a DNA sequence may vary from that shown in SEQ ID NO:1, SEQ ID NO:7, and SEQ ID NO:9 and still encode a LERK-6 protein having the amino acid sequence of SEQ ID NO:2, SEQ ID NO:8, or SEQ ID NO:10, respectively. Such variant DNA sequences may result from silent mutations (e.g., occurring during PCR amplification), or may be the product of deliberate mutagenesis of a native sequence.

The invention provides equivalent isolated DNA sequences encoding biologically active LERK-6, selected from: (a) cDNA comprising the nucleotide sequence presented in SEQ ID NO:1, SEQ ID NO:7, or SEQ ID NO:9; (b) DNA capable of hybridization to a DNA of (a) under moderately stringent conditions and that encodes biologically active LERK-6; and (c) DNA that is degenerate as a result of the genetic code to a DNA defined in (a), or (b) and that encodes biologically active LERK-6. LERK-6 proteins encoded by such DNA equivalent sequences are encompassed by the invention.

DNAs that are equivalents to the DNA sequence of SEQ ID NO:1, and SEQ ID NO:7 or SEQ ID NO:9 will hybridize under moderately and severely stringent conditions to DNA sequences that encode polypeptides comprising amino acids 1-184 of SEQ ID NO:2, SEQ ID NO:8, or amino acids 1-184 of SEQ ID NO:10. Examples of LERK-6 proteins encoded by such DNA, include, but are not limited to, LERK-6 fragments (soluble or GPI-linked) and LERK-6 proteins comprising inactivated N-glycosylation site(s), inactivated KEX2 protease processing site(s), or conservative amino acid substitution(s), as described above. LERK-6 proteins encoded by DNA derived from

other mammalian species, wherein the DNA will hybridize to the cDNA of SEQ ID NO:1, SEQ ID NO:7 or SEQ ID NO:9 are also encompassed by the present invention.

5 Variants possessing the requisite ability to bind hek/elk receptor may be identified by any suitable assay. Biological activity of LERK-6 may be determined, for example, by competition for binding to the ligand binding domain of hek/elk receptor (i.e. competitive binding assays).

10 One type of a competitive binding assay for a LERK-6 polypeptide uses a radiolabeled, soluble LERK-6 and intact hek/elk-expressing cells. Instead of intact cells, one could substitute soluble hek/elk:Fc fusion proteins (such as a hek:Fc or elk:Fc fusion protein) bound to a solid phase through the interaction of a Protein A, Protein G or an antibody to the hek, elk or Fc portions of the molecule, with the Fc region of the fusion protein. Another type of competitive binding assay utilizes radiolabeled soluble hek or
15 elk receptors such as a hek:Fc or elk:Fc fusion protein, and intact cells expressing LERK-6.

Competitive binding assays can be performed following conventional methodology. For example, radiolabeled LERK-6 can be used to compete with a putative
20 LERK-6 homolog to assay for binding activity against surface-bound hek/elk. Qualitative results can be obtained by competitive autoradiographic plate binding assays, or Scatchard plots may be utilized to generate quantitative results.

Alternatively, hek/elk-binding proteins, such as LERK-6 and anti-hek/elk
25 antibodies, can be bound to a solid phase such as a column chromatography matrix or a similar substrate suitable for identifying, separating or purifying cells that express hek/elk on their surface. Binding of a hek/elk-binding protein to a solid phase contacting surface can be accomplished by any means, for example, by constructing a LERK-6:Fc fusion protein and binding such to the solid phase through the interaction of Protein A or Protein
30 G. Various other means for fixing proteins to a solid phase are well known in the art and are suitable for use in the present invention. For example, magnetic microspheres can be coated with LERK-6 and held in the incubation vessel through a magnetic field. Suspensions of cell mixtures containing hek/elk-expressing cells are contacted with the solid phase that has LERK-6 polypeptides thereon. Cells having hek/elk on their surface
35 bind to the fixed LERK-6 and unbound cells then are washed away. This affinity-binding method is useful for purifying, screening or separating such hek/elk-expressing cells from solution. Methods of releasing positively selected cells from the solid phase are known in

the art and encompass, for example, the use of enzymes. Such enzymes are preferably non-toxic and non-injurious to the cells and are preferably directed to cleaving the cell-surface binding partner. In the case of hek/elk:LERK-6 interactions, the enzyme preferably would cleave the hek/elk receptor, thereby freeing the resulting cell suspension from the "foreign" LERK-6 material. The purified cell population, especially if obtained from fetal tissue, then may be used to repopulate mature (adult) tissues. For example, such purified neural cells can be administered to a patient having a neurodegenerative disorder such as Parkinson's disease, Alzheimer's disease, Lou Gehrig's disease, etc.

Alternatively, mixtures of cells suspected of containing hek/elk⁺ cells first can be incubated with biotinylated LERK-6. Incubation periods are typically at least one hour in duration to ensure sufficient binding to hek/elk. The resulting mixture then is passed through a column packed with avidin-coated beads, whereby the high affinity of biotin for avidin provides the binding of the cell to the beads. Use of avidin-coated beads is known in the art. See Berenson, et al. *J. Cell. Biochem.*, 10D:239 (1986). Wash of unbound material and the release of the bound cells is performed using conventional methods.

As described above, LERK-6 can be used to separate cells expressing hek/elk. In an alternative method, LERK-6 or an extracellular domain or a fragment thereof can be conjugated to a detectable moiety such as ¹²⁵I to detect hek/elk-expressing cells. Radiolabeling with ¹²⁵I can be performed by any of several standard methodologies that yield a functional ¹²⁵I-LERK-6 molecule labeled to high specific activity. Or an iodinated or biotinylated antibody against the hek/elk region or the Fc region of the molecule could be used. Another detectable moiety such as an enzyme that can catalyze a colorimetric or fluorometric reaction, biotin or avidin may be used. Cells to be tested for hek/elk-expression can be contacted with labeled LERK-6. After incubation, unbound labeled LERK-6 is removed and binding is measured using the detectable moiety.

The binding characteristics of LERK-6 (including variants) may also be determined using the conjugated, soluble hek/elk (for example, ¹²⁵I-hek/elk:Fc) in competition assays similar to those described above. In this case, however, intact cells expressing hek/elk bound to a solid substrate, are used to measure the extent to which a sample containing a putative LERK-6 variant competes for binding with a conjugated a soluble hek/elk to LERK-6.

Other means of assaying for LERK-6 include the use of anti-LERK-6 antibodies, cell lines that proliferate in response to LERK-6, or recombinant cell lines that express hek/elk and proliferate in the presence of LERK-6.

5 The LERK-6 proteins disclosed herein also may be employed to measure the biological activity of elk or hek proteins in terms of their binding affinity for LERK-6. As one example, LERK-6 may be used in determining whether biological activity is retained after modification of an elk or hek protein (e.g., chemical modification, truncation, mutation, etc.). The biological activity of an elk or hek protein thus can be
10 ascertained before it is used in a research study, or possibly in the clinic, for example.

 LERK-6 proteins find use as reagents that may be employed by those conducting "quality assurance" studies, e.g., to monitor shelf life and stability of elk protein under different conditions. To illustrate, LERK-6 may be employed in a binding affinity study
15 to measure the biological activity of an elk protein that has been stored at different temperatures, or produced in different cell types. The binding affinity of the modified elk protein for LERK-6 is compared to that of an unmodified elk protein to detect any adverse impact of the modifications on biological activity of elk. Likewise, the biological activity of a hek protein can be assessed using LERK-6.

20 LERK-6 polypeptides also find use as carriers for delivering agents attached thereto to cells bearing the elk or hek cell surface receptor. Expression of hek antigen has been reported for certain leukemic cell lines, including the human T-cell leukemia cell line designated JM and the human pre-B cell leukemia cell line designated LK63 (Boyd et al., *J. Biol. Chem.* 267:3262, 1992, and Wicks et al., *Proc. Natl. Acad. Sci. USA*, 89:1611, 1992). LERK-6 proteins thus can be used to deliver diagnostic or therapeutic agents to
25 these cells (or to other cell types found to express hek on the cell surface) in *in vitro* or *in vivo* procedures.

30 One example of such use is to expose a hek⁺ or elk⁺ leukemic cell line to a therapeutic agent/LERK-6 conjugate to assess whether the agent exhibits cytotoxicity toward the leukemic cells. A number of different therapeutic agents attached to LERK-6 may be included in an assay to detect and compare the cytotoxic effect of the agents on the leukemic cells. LERK-6/diagnostic agent conjugates may be employed to detect the
35 presence of hek⁺ cells *in vitro* or *in vivo*.

Diagnostic and therapeutic agents that may be attached to a LERK-6 polypeptide include, but are not limited to, drugs, toxins, radionuclides, chromophores, enzymes that catalyze a colorimetric or fluorometric reaction, and the like, with the particular agent being chosen according to the intended application. Examples of drugs include those used in treating various forms of cancer, e.g., nitrogen mustards such as L-phenylalanine nitrogen mustard or cyclophosphamide, intercalating agents such as cis-diaminodichloroplatinum, antimetabolites such as 5-fluorouracil, vinca alkaloids such as vincristine, and antibiotics such as bleomycin, doxorubicin, daunorubicin, and derivatives thereof. Among the toxins are ricin, abrin, diphtheria toxin, *Pseudomonas aeruginosa* exotoxin A, ribosomal inactivating proteins, mycotoxins such as trichothecenes, and derivatives and fragments (e.g., single chains) thereof. Radionuclides suitable for diagnostic use include, but are not limited to, ^{123}I , ^{131}I , $^{99\text{m}}\text{Tc}$, ^{111}In , and ^{76}Br . Radionuclides suitable for therapeutic use include, but are not limited to, ^{131}I , ^{211}At , ^{77}Br , ^{186}Re , ^{188}Re , ^{212}Pb , ^{212}Bi , ^{109}Pd , ^{64}Cu , and ^{67}Cu .

Such agents may be attached to the LERK-6 by any suitable conventional procedure. LERK-6, being a protein, comprises functional groups on amino acid side chains that can be reacted with functional groups on a desired agent to form covalent bonds, for example. Alternatively, the protein or agent may be derivatized to generate or attach a desired reactive functional group. The derivatization may involve attachment of one of the bifunctional coupling reagents available for attaching various molecules to proteins (Pierce Chemical Company, Rockford, Illinois). A number of techniques for radiolabeling proteins are known. Radionuclide metals may be attached to LERK-6 by using a suitable bifunctional chelating agent, for example.

Conjugates comprising LERK-6 and a suitable diagnostic or therapeutic agent (preferably covalently linked) are thus prepared. The conjugates are administered or otherwise employed in an amount appropriate for the particular application.

Another use of the LERK-6 of the present invention is as a research tool for studying the role that LERK-6, in conjunction with elk or hek, may play in growth or differentiation of cells bearing the elk or hek receptor. The LERK-6 polypeptides of the present invention also may be employed in *in vitro* assays for detection of elk or LERK-6 or the interactions thereof. Likewise, LERK-6 finds use in assays for hek or the interaction of LERK-6 with hek.

As discussed above, when various rat tissues were analyzed for elk mRNA, transcripts were detected only in brain and testis (Lhotak et al., *supra*). Binding of LERK-6 to eph-related receptors on neural tissue is believed to exert a neuroprotective or neurotrophic effect.

5

LERK-6 finds use as a tissue culture reagent. A LERK-6 protein can be added to neurons cultured *in vitro* to enhance the viability or prolong the lifespan of the cultured neurons, thus facilitating research studies, and possible clinical treatment of neural tissue.

10

One embodiment of the present invention is directed to a method of treating disorders of neural tissue, involving contacting the neural tissue with LERK-6. Such disorders include injury or neurologic diseases, either chronic or acute. A LERK-6 protein may be administered to a mammal to treat such an injury or disease. In one embodiment of the invention, LERK-6 is employed in treating neurodegenerative conditions characterized or mediated, at least in part, by the mechanism of neural death known as excitotoxicity. In addition, LERK-6 may be administered to a mammal to exert a trophic effect on neural tissue. In a patient suffering loss of or damage to neurons due to injury or disease, LERK-6 may enhance the viability of those neurons that have survived.

20

LERK-6 polypeptides of the invention can be formulated according to known methods used to prepare pharmaceutically useful compositions. LERK-6 can be combined in admixture, either as the sole active material or with other known active materials, with pharmaceutically suitable diluents (e.g., Tris-HCl, acetate, phosphate), preservatives (e.g., Thimerosal, benzyl alcohol, parabens), emulsifiers, solubilizers, adjuvants and/or carriers. Suitable carriers and their formulations are described in Remington's Pharmaceutical Sciences, 16th ed. 1980, Mack Publishing Co. In addition, such compositions can contain LERK-6 complexed with polyethylene glycol (PEG), metal ions, or incorporated into polymeric compounds such as polyacetic acid, polyglycolic acid, hydrogels, etc., or incorporated into liposomes, microemulsions, micelles, unilamellar or multilamellar vesicles, erythrocyte ghosts or spheroblasts. Such compositions will influence the physical state, solubility, stability, rate of *in vivo* release, and rate of *in vivo* clearance of LERK-6. LERK-6 can also be conjugated to antibodies against tissue-specific receptors, ligands or antigens, or coupled to ligands of tissue-specific receptors. Where hek/elk is found on neoplastic cells, the LERK-6 may be conjugated to a toxin whereby LERK-6 is used to deliver the toxin to the specific site, or

30

may be used to sensitize such neoplastic cells to subsequently administered anti-neoplastic agents.

LERK-6 can be administered topically, parenterally, or by inhalation. The term
5 "parenteral" includes subcutaneous injections, intravenous, intramuscular, intracisternal
injection, or infusion techniques. These compositions will typically contain an effective
amount of the LERK-6, alone or in combination with an effective amount of any other
active material. Such dosages and desired drug concentrations contained in the
compositions may vary depending upon many factors, including the intended use,
10 patient's body weight and age, and route of administration. Preliminary doses can be
determined according to animal tests, and the scaling of dosages for human
administration can be performed according to art-accepted practices.

LERK-6 polypeptides may exist as oligomers, such as covalently-linked or non-
15 covalently-linked dimers or trimers. Oligomers may be linked by disulfide bonds formed
between cysteine residues on different LERK-6 polypeptides. In one embodiment of the
invention, a LERK-6 dimer is created by fusing LERK-6 to the Fc region of an antibody
(e.g., IgG1) in a manner that does not interfere with binding of LERK-6 to the hek/elk
ligand-binding domain. The Fc polypeptide preferably is fused to the C-terminus of a
20 soluble LERK-6 (comprising only the receptor-binding). General preparation of fusion
proteins comprising heterologous polypeptides fused to various portions of antibody-
derived polypeptides (including the Fc domain) has been described, e.g., by Ashkenazi et
al. (*PNAS USA* 88:10535, 1991) and Byrn et al. (*Nature* 344:677, 1990), hereby
incorporated by reference. A gene fusion encoding the LERK-6:Fc fusion protein is
25 inserted into an appropriate expression vector. LERK-6:Fc fusion proteins are allowed to
assemble much like antibody molecules, whereupon interchain disulfide bonds form
between Fc polypeptides, yielding divalent LERK-6. If fusion proteins are made with
both heavy and light chains of an antibody, it is possible to form a LERK-6 oligomer with
as many as four LERK-6 extracellular regions. Alternatively, one can link two soluble
30 LERK-6 domains with a peptide linker.

Suitable host cells for expression of LERK-6 polypeptides include prokaryotes,
yeast or higher eukaryotic cells. Appropriate cloning and expression vectors for use with
bacterial, fungal, yeast, and mammalian cellular hosts are described, for example, in
35 Pouwels et al. *Cloning Vectors: A Laboratory Manual*, Elsevier, New York, (1985).
Cell-free translation systems could also be employed to produce LERK-6 polypeptides
using RNAs derived from DNA constructs disclosed herein.

Prokaryotes include gram negative or gram positive organisms, for example, *E. coli* or *Bacilli*. Suitable prokaryotic host cells for transformation include, for example, *E. coli*, *Bacillus subtilis*, *Salmonella typhimurium*, and various other species within the genera *Pseudomonas*, *Streptomyces*, and *Staphylococcus*. In a prokaryotic host cell, such as *E. coli*, a LERK-6 polypeptide may include an N-terminal methionine residue to facilitate expression of the recombinant polypeptide in the prokaryotic host cell. The N-terminal Met may be cleaved from the expressed recombinant LERK-6 polypeptide.

LERK-6 polypeptides may be expressed in yeast host cells, preferably from the *Saccharomyces* genus (e.g., *S. cerevisiae*). Other genera of yeast, such as *Pichia*, *K. lactis* or *Kluyveromyces*, may also be employed. Yeast vectors will often contain an origin of replication sequence from a 2 μ yeast plasmid, an autonomously replicating sequence (ARS), a promoter region, sequences for polyadenylation, sequences for transcription termination, and a selectable marker gene. Suitable promoter sequences for yeast vectors include, among others, promoters for metallothionein, 3-phosphoglycerate kinase (Hitzeman et al., *J. Biol. Chem.* 255:2073, 1980) or other glycolytic enzymes (Hess et al., *J. Adv. Enzyme Reg.* 7:149, 1968; and Holland et al., *Biochem.* 17:4900, 1978), such as enolase, glyceraldehyde-3-phosphate dehydrogenase, hexokinase, pyruvate decarboxylase, phosphofructokinase, glucose-6-phosphate isomerase, 3-phosphoglycerate mutase, pyruvate kinase, triosephosphate isomerase, phosphoglucose isomerase, and glucokinase. Other suitable vectors and promoters for use in yeast expression are further described in Hitzeman, EPA-73,657 or in Fleer et al., *Gene*, 107:285-195 (1991); and van den Berg et al., *Bio/Technology*, 8:135-139 (1990). Another alternative is the glucose-repressible ADH2 promoter described by Russell et al. (*J. Biol. Chem.* 258:2674, 1982) and Beier et al. (*Nature* 300:724, 1982). Shuttle vectors replicable in both yeast and *E. coli* may be constructed by inserting DNA sequences from pBR322 for selection and replication in *E. coli* (Amp^r gene and origin of replication) into the above-described yeast vectors.

The yeast α -factor leader sequence may be employed to direct secretion of the LERK-6 polypeptide. The α -factor leader sequence is often inserted between the promoter sequence and the structural gene sequence. See, e.g., Kurjan et al., *Cell* 30:933, 1982; Bitter et al., *Proc. Natl. Acad. Sci. USA* 81:5330, 1984; U. S. Patent 4,546,082; and EP 324,274. Other leader sequences suitable for facilitating secretion of recombinant polypeptides from yeast hosts are known to those of skill in the art. A leader sequence

may be modified near its 3' end to contain one or more restriction sites. This will facilitate fusion of the leader sequence to the structural gene.

5 Yeast transformation protocols are known to those of skill in the art. One such protocol is described by Hinnen et al., *Proc. Natl. Acad. Sci. USA* 75:1929, 1978. The Hinnen et al. protocol selects for Trp⁺ transformants in a selective medium, wherein the selective medium consists of 0.67% yeast nitrogen base, 0.5% casamino acids, 2% glucose, 10 µg/ml adenine and 20 µg/ml uracil.

10 Yeast host cells transformed by vectors containing ADH2 promoter sequence may be grown for inducing expression in a "rich" medium. An example of a rich medium is one consisting of 1% yeast extract, 2% peptone, and 1% glucose supplemented with 80 µg/ml adenine and 80 µg/ml uracil. Derepression of the ADH2 promoter occurs when glucose is exhausted from the medium.

15 Mammalian or insect host cell culture systems could also be employed to express recombinant LERK-6 polypeptides. Baculovirus systems for production of heterologous proteins in insect cells are reviewed by Luckow and Summers, *Bio/Technology* 6:47 (1988). Established cell lines of mammalian origin also may be employed. Examples of
20 suitable mammalian host cell lines include the COS-7 line of monkey kidney cells (ATCC CRL 1651) (Gluzman et al., *Cell* 23:175, 1981), L cells, C127 cells, 3T3 cells (ATCC CCL 163), Chinese hamster ovary (CHO) cells, HeLa cells, and BHK (ATCC CRL 10) cell lines, and the CV-1/EBNA-1 cell line derived from the African green monkey kidney cell line CV1 (ATCC CCL 70) as described by McMahan et al. (*EMBO J.*
25 10: 2821, 1991).

Transcriptional and translational control sequences for mammalian host cell expression vectors may be excised from viral genomes. Commonly used promoter sequences and enhancer sequences are derived from Polyoma virus, Adenovirus 2,
30 Simian Virus 40 (SV40), and human cytomegalovirus. DNA sequences derived from the SV40 viral genome, for example, SV40 origin, early and late promoter, enhancer, splice, and polyadenylation sites may be used to provide other genetic elements for expression of a structural gene sequence in a mammalian host cell. Viral early and late promoters are particularly useful because both are easily obtained from a viral genome as a fragment
35 which may also contain a viral origin of replication (Fiers et al., *Nature* 273:113, 1978). Smaller or larger SV40 fragments may also be used, provided the approximately 250 bp

sequence extending from the *Hind* III site toward the *Bgl* I site located in the SV40 viral origin of replication site is included.

Exemplary expression vectors for use in mammalian host cells can be constructed
5 as disclosed by Okayama and Berg (*Mol. Cell. Biol.* 3:280, 1983). A useful system for
stable high level expression of mammalian cDNAs in C127 murine mammary epithelial
cells can be constructed substantially as described by Cosman et al. (*Mol. Immunol.*
23:935, 1986). A useful high expression vector, PMLSV N1/N4, described by Cosman et
al., *Nature* 312:768, 1984 has been deposited as ATCC 39890. Additional useful
10 mammalian expression vectors are described in EP-A-0367566, and in U.S. Patent
Application Serial No. 07/701,415, filed May 16, 1991, incorporated by reference herein.
The vectors may be derived from retroviruses. In place of the native signal sequence, and
in addition to an initiator methionine, a heterologous signal sequence may be added, such
as the signal sequence for IL-7 described in United States Patent 4,965,195; the signal
15 sequence for IL-2 receptor described in Cosman et al., *Nature* 312:768 (1984); the IL-4
signal peptide described in EP 367,566; the type I IL-1 receptor signal peptide described
in U.S. Patent 4,968,607; and the type II IL-1 receptor signal peptide described in EP
460,846.

20 LERK-6 as an isolated, purified or homogeneous protein according to the
invention may be produced by recombinant expression systems as described above or
purified from naturally occurring cells. LERK-6 can be purified to substantial
homogeneity, as indicated by a single protein band upon analysis by SDS-polyacrylamide
gel electrophoresis (SDS-PAGE).

25 One process for producing LERK-6 comprises culturing a host cell transformed
with an expression vector comprising a DNA sequence that encodes LERK-6 under
conditions sufficient to promote expression of LERK-6. LERK-6 is then recovered from
culture medium or cell extracts, depending upon the expression system employed. As is
30 known to the skilled artisan, procedures for purifying a recombinant protein will vary
according to such factors as the type of host cells employed and whether or not the
recombinant protein is secreted into the culture medium.

For example, when expression systems that secrete the recombinant protein are
35 employed, the culture medium first may be concentrated using a commercially available
protein concentration filter, for example, an Amicon or Millipore Pellicon ultrafiltration
unit. Following the concentration step, the concentrate can be applied to a purification

matrix such as a gel filtration medium. Alternatively, an anion exchange resin can be employed, for example, a matrix or substrate having pendant diethylaminoethyl (DEAE) groups. The matrices can be acrylamide, agarose, dextran, cellulose or other types commonly employed in protein purification. Alternatively, a cation exchange step can be employed. Suitable cation exchangers include various insoluble matrices comprising sulfopropyl or carboxymethyl groups. Sulfopropyl groups are preferred. Finally, one or more reversed-phase high performance liquid chromatography (RP-HPLC) steps employing hydrophobic RP-HPLC media, (e.g., silica gel having pendant methyl or other aliphatic groups) can be employed to further purify LERK-6. Some or all of the foregoing purification steps, in various combinations, are well known and can be employed to provide a substantially homogeneous recombinant protein.

It is possible to utilize an affinity column comprising the ligand-binding domain of hek/elk receptors to affinity-purify expressed LERK-6 polypeptides. LERK-6 polypeptides can be removed from an affinity column using conventional techniques, e.g., in a high salt elution buffer and then dialyzed into a lower salt buffer for use or by changing pH or other components depending on the affinity matrix utilized. Alternatively, the affinity column may comprise an antibody that binds LERK-6. Example 5 describes a procedure for employing LERK-6 of the invention to generate monoclonal antibodies directed against LERK-6.

Recombinant protein produced in bacterial culture can be isolated by initial disruption of the host cells, centrifugation, extraction from cell pellets if an insoluble polypeptide, or from the supernatant fluid if a soluble polypeptide, followed by one or more concentration, salting-out, ion exchange, affinity purification or size exclusion chromatography steps. Finally, RP-HPLC can be employed for final purification steps. Microbial cells can be disrupted by any convenient method, including freeze-thaw cycling, sonication, mechanical disruption, or use of cell lysing agents.

Transformed yeast host cells are preferably employed to express LERK-6 as a secreted polypeptide in order to simplify purification. Secreted recombinant polypeptide from a yeast host cell fermentation can be purified by methods analogous to those disclosed by Urdal et al. (*J. Chromatog.* 296:171, 1984). Urdal et al. describe two sequential, reversed-phase HPLC steps for purification of recombinant human IL-2 on a preparative HPLC column.

Useful fragments of the LERK-6 nucleic acids include antisense or sense oligonucleotides comprising a single-stranded nucleic acid sequence (either RNA or

DNA) capable of binding to target LERK-6 mRNA (sense) or LERK-6 DNA (antisense) sequences. Antisense or sense oligonucleotides, according to the present invention, comprise a fragment of the coding region of LERK-6 cDNA. Such a fragment generally comprises at least about 14 nucleotides, preferably from about 14 to about 30 nucleotides.

- 5 The ability to derive an antisense or a sense oligonucleotide, based upon a cDNA sequence encoding a given protein is described in, for example, Stein and Cohen (*Cancer Res.* 48:2659, 1988) and van der Krol et al. (*BioTechniques* 6:958, 1988).

Binding of antisense or sense oligonucleotides to target nucleic acid sequences results in the formation of duplexes that block transcription or translation of the target
10 sequence by one of several means, including enhanced degradation of the duplexes, premature termination of transcription or translation, or by other means. The antisense oligonucleotides thus may be used to block expression of LERK-6 proteins. Antisense or sense oligonucleotides further comprise oligonucleotides having modified sugar-phosphodiester backbones (or other sugar linkages, such as those described in
15 WO91/06629) and wherein such sugar linkages are resistant to endogenous nucleases. Such oligonucleotides with resistant sugar linkages are stable *in vivo* (i.e., capable of resisting enzymatic degradation) but retain sequence specificity to be able to bind to target nucleotide sequences. Other examples of sense or antisense oligonucleotides include those oligonucleotides which are covalently linked to organic moieties, such as
20 those described in WO 90/10448, and other moieties that increases affinity of the oligonucleotide for a target nucleic acid sequence, such as poly-(L-lysine). Further still, intercalating agents, such as ellipticine, and alkylating agents or metal complexes may be attached to sense or antisense oligonucleotides to modify binding specificities of the antisense or sense oligonucleotide for the target nucleotide sequence.

25 Antisense or sense oligonucleotides may be introduced into a cell containing the target nucleic acid sequence by any gene transfer method, including, for example, CaPO₄-mediated DNA transfection, electroporation, or by using gene transfer vectors such as Epstein-Barr virus. Antisense or sense oligonucleotides are preferably introduced into a cell containing the target nucleic acid sequence by insertion of the antisense or sense
30 oligonucleotide into a suitable retroviral vector, then contacting the cell with the retrovirus vector containing the inserted sequence, either *in vivo* or *ex vivo*. Suitable retroviral vectors include, but are not limited to, those derived from the murine retrovirus M-MuLV, N2 (a retrovirus derived from M-MuLV), or the double copy vectors designated DCT5A, DCT5B and DCT5C (see PCT Application US 90/02656).

Sense or antisense oligonucleotides also may be introduced into a cell containing the target nucleotide sequence by formation of a conjugate with a ligand binding molecule, as described in WO 91/04753. Suitable ligand binding molecules include, but are not limited to, cell surface receptors, growth factors, other cytokines, or other ligands that bind to cell surface receptors. Preferably, conjugation of the ligand binding molecule does not substantially interfere with the ability of the ligand binding molecule to bind to its corresponding molecule or receptor, or block entry of the sense or antisense oligonucleotide or its conjugated version into the cell.

Alternatively, a sense or an antisense oligonucleotide may be introduced into a cell containing the target nucleic acid sequence by formation of an oligonucleotide-lipid complex, as described in WO 90/10448. The sense or antisense oligonucleotide-lipid complex is preferably dissociated within the cell by an endogenous lipase.

In addition to the above, the following examples are provided to illustrate particular embodiments and not to limit the scope of the invention.

EXAMPLE 1

Cloning of Murine LERK-6 cDNA

This example describes a procedure for isolating a DNA sequence encoding murine LERK-6 of the invention. A commercially available 11.5 day murine embryonic cDNA library was obtained from Clonotech Laboratories, Inc., Palo Alto, California. The library was plated according to the procedures detailed in the manual provided by Clonotech, and screened using the ³²P-radiolabelled probes. The probes each were generated using standard techniques. Generally, polymerase chain reaction (PCR) (Mullis and Faloona, *Meth. Enzymol.* 155:335-350, 1987) amplifications were performed using two sets of primers. The first set,

GATATTTACT GCCCGCACTA CAACAGCT

SEQ ID NO:3

AGAGAAGGCG CTGTAGCGCT GGAAC

SEQ ID NO:4

30

was used to generate amplified double stranded DNA fragments from the DNA of LERK-3. LERK-3 is the subject of copending patent application Serial Nos. 08/109,745; 08/114,426, 08/161,132 and 08/240,124 filed May 9, 1994. The probe from LERK-3 comprised nucleotides 260 through 481 of the SEQ ID NO:1 of copending application serial no. 08/240,124 filed May 9, 1994. The second set of primers,

35

ACGTAGTCTA CTGGAAGTCC AGTAACCCCA G

SEQ ID NO:5

AGCCTCAAGC ACTGGCCAGA ACTCTCTCTG GAGT SEQ ID NO:6

was used to generate amplified double stranded DNA fragments from the DNA of LERK-4. LERK-4 also is the subject of copending patent application Serial Nos. 08/109,745; 5 08/114,426, 08/161,132 and 08/240,124 filed May 9, 1994. The probe from LERK-3 comprised nucleotides 110 through 467 of the SEQ ID NO:3 of copending application serial no. 08/240,124 filed May 9, 1994. The fragments were radiolabelled with ³²P, and used as probes to screen a murine embryonic library.

10 Screening with the probes was conducted by conventional procedures, and hybridizing clones were identified. The hybridizing conditions consisted of 42°C and 50% Starks washed to 0.1X SSC at 63°C. The nucleotide sequence of the cDNA insert of one individual clone isolated from the murine embryonic library, designated clone λ13, was determined. A substantially complete cDNA sequence of the coding region of the 15 clone λ13 cDNA, and the amino acid sequence encoded thereby, are presented in SEQ ID NO:1 and SEQ ID NO:2, respectively. The open-reading frame within this sequence encodes a protein of 184 amino acids. Since the N-termini of the proteins of the LERK family are non-homologous, it is believed that the first amino acid of SEQ ID NO:1 (Ala) is located at or very near the native N-terminus. In addition, because LERK-6 of this 20 invention is homologous with other biologically active LERK molecules, there is a significant probability that the cDNA of SEQ ID NO:1 encodes a biologically active protein. Secretion of such active protein is possible through the procedures described above with respect to addition of a suitable initiator methionine and a suitable signal sequence, such as that for murine IL-7. The amino acid sequence of LERK-6 is 58% 25 homologous to that of a cytokine termed LERK-3.

A cell lysate containing clone λ13 DNA (the LERK-6 cDNA in λgt10) was deposited with the American Type Culture Collection, Rockville, MD, USA on July 15, 1994, and assigned accession number ATCC 75829. The deposit was made under the 30 terms of the Budapest Treaty.

EXAMPLE 2

Cloning of Human LERK-6

35 This example describes the cloning of nucleic acids encoding human LERK-6. Human LERK-6 sequence is from a genomic clone isolated by probing a human genomic library with a murine LERK-6 cDNA probe encoding amino acids 30-173. Plaque containing filters were hybridized to ³²P-labeled probe at 55°C for 24 h, washed to 0.5 x

SSC at 55°C, and exposed to X-ray film (Kodak XAR-5). A sequence of nucleic acids contained in such clone is shown below, also in SEQ ID NO:7, and the amino acids encoded thereby are shown in SEQ ID NO:8:

```

5      G TTC CAC GCA GGC GCG GGG GAC GAC GGC GGG GGC TAC ACG GTG GAG      46
      Phe His Ala Gly Ala Gly Asp Asp Gly Gly Gly Tyr Thr Val Glu
        1          5          10          15

10     GTG AGC ATC AAT GAC TAC CTG GAC ATC TAC TGC CCG CAC TAT GGG GCG      94
      Val Ser Ile Asn Asp Tyr Leu Asp Ile Tyr Cys Pro His Tyr Gly Ala
                20          25          30

15     CCG CTG CCG CCG GCC GAG CGC ATG GAG CAC TAC GTG CTG TAC ATG GTC      142
      Pro Leu Pro Pro Ala Glu Arg Met Glu His Tyr Val Leu Tyr Met Val
                35          40          45

20     AAC GGC GAG GGC CAC GCC TCC TGC GAC CAC CGC CAG CGC GGC TTC AAG      190
      Asn Gly Glu Gly His Ala Ser Cys Asp His Arg Gln Arg Gly Phe Lys
                50          55          60

25     CGC TGG GAG TGC AAC CGG CCC GCG GCG CCC GGG GGG CCG CTC AAG TTC      238
      Arg Trp Glu Cys Asn Arg Pro Ala Ala Pro Gly Gly Pro Leu Lys Phe
                65          70          75

30     TCG GAG AAG TTC CAG CTC TTC ACG CCC TTC TCC CTG GGC TTC GAG TTC      286
      Ser Glu Lys Phe Gln Leu Phe Thr Pro Phe Ser Leu Gly Phe Glu Phe
        80          85          90          95

30     CGG CCC GGC CAC GAG TAT TAC TAC ATC T      314
      Arg Pro Gly His Glu Tyr Tyr Tyr Ile
                100

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35 Compared to murine LERK-6, human LERK-6 possesses a 98.077 percent similarity and 93.269 percent identity.

A complete human LERK-6 DNA was isolated by probing a human genomic DNA library (Stratagene, La Jolla, CA) with the nucleic acid sequence shown in SEQ ID NO:7. Plaque containing filters were hybridized to the ³²P-labeled probe at 55°C for 24 h, washed to 0.5 x SSC at 55°C, and exposed to X-ray film. Two different clones were isolated using this technique. Then the same isolation process was repeated using the human genomic DNA library from a different preparation (Stratagene, La Jolla, CA). This process resulted in the isolation of a third clone having a different nucleotide sequence. The three isolated clones were restriction mapped and then subcloned into bluescript plasmid and sequenced. Using the murine LERK-6 sequence as a template the introns and exons of the clones were identified. The full length human LERK-6 encoding DNA sequence as determined by comparing the coding regions of the murine LERK-6 is identified in SEQ ID NO:9 and the amino acid sequence encoded by the full length DNA sequence is identified in SEQ ID NO:10.

Plasmids including the cloned LERK-6 DNA were deposited with the American Type Culture Collection, Rockville, MD, USA on August __ 1997, and assigned accession number ATCC _____. The deposit was made under the terms of the Budapest Treaty.

EXAMPLE 3

Preparation of Soluble elk:Fc Fusion Protein

This example describes construction of an expression vector encoding a soluble elk:Fc fusion protein. This fusion protein can be employed in the binding assays of Example 5, to determine the binding characteristics of LERK-6.

A DNA and encoded amino acid sequence for rat elk cDNA is presented in Lhotak et al. (*Mol. Cell. Biol.* 11:2496, 1991), hereby incorporated by reference. The rat elk protein has a 538 amino acid extracellular domain, a 25 amino acid transmembrane domain, and a 419 amino acid cytoplasmic domain.

A rat elk cDNA fragment was fused to the 5' end of cDNA encoding the Fc portion of a human IgG1 antibody. The rat elk cDNA can be obtained from T. Pawson (Samuel Lunenfeld Research Institute, Mt. Sinai Hospital, Toronto). An Asp718 restriction endonuclease cleavage site was introduced upstream of the elk coding region. An Asp 718-BglII fragment of rat elk cDNA (comprising the entire extracellular domain, the transmembrane region, and a small portion of the cytoplasmic domain) was isolated.

DNA encoding a single polypeptide chain comprising the Fc region of a human IgG1 antibody was cloned into the *SpeI* site of the pBLUESCRIPT SK® vector, which is commercially available from Stratagene Cloning Systems, La Jolla, California. This plasmid vector is replicable in *E. coli* and contains a polylinker segment that includes 21 unique restriction sites. The nucleotide sequence of the cloned DNA, along with the amino acid sequence of the Fc polypeptide encoded thereby, are described in PCT application WO 93/10151, hereby incorporated by reference. A unique *BglII* site has been introduced, and encompasses the codons for amino acids three and four of the Fc polypeptide. The encoded Fc polypeptide extends from the N-terminal hinge region to the native C-terminus, i.e., is an essentially full-length antibody Fc region.

The above-described Asp718-BglII elk cDNA fragment was cloned into the pBLUESCRIPT SK® vector containing the Fc cDNA, such that the elk cDNA is positioned upstream of the Fc cDNA. Single stranded DNA derived from the resulting

gene fusion was mutagenized by the method described in Kunkel (*Proc. Natl. Acad. Sci. USA* 82:488, 1985) and Kunkel et al. (*Methods in Enzymol.* 154:367, 1987) in order to perfectly fuse the entire extracellular domain of elk to the Fc sequence. The mutagenized DNA was sequenced to confirm that the proper nucleotides had been removed (i.e., that
5 the transmembrane region and partial cytoplasmic domain DNA were deleted) and that the elk and Fc sequences were in the same reading frame.

The elk:Fc fusion protein preferably is synthesized in mammalian host cells, such as CV1-EBNA or COS-7 cells. The elk:Fc gene fusion was excised and inserted into a mammalian expression vector designated HAV-EO (Dower et al., *J. Immunol.* 142:4314,
10 1989). Mammalian host cells were transfected with the resulting recombinant expression vector and cultivated to allow transient expression of the fusion protein, which was secreted into the culture medium *via* the elk signal peptide. The elk:Fc fusion protein was purified by affinity chromatography, using a protein A sepharose column.

15

EXAMPLE 4

Preparation of Soluble hek:Fc Fusion Protein

This example describes construction of an expression vector encoding a soluble hek:Fc fusion protein. This fusion protein can be employed in the binding assays of Example 5, to determine the binding characteristics of LERK-6.

20

A DNA and encoded amino acid sequence for human hek cDNA is presented in Wicks et al. (*Proc. Nat'l. Acad. Sci. USA*, 89:1611, 1992), incorporated herein by reference. This hek protein comprises (from N- to C-terminus) an extracellular domain, a transmembrane domain, and a cytoplasmic domain.

Two DNA fragments, one encoding an N-terminal fragment of the extracellular
25 domain of hek and the other encoding a C-terminal fragment of the hek extracellular domain, were isolated by polymerase chain reactions (PCR) conducted under standard conditions, using oligonucleotide primers based on the hek nucleotide sequence published by Wicks et al., *supra*. The template for the PCR was cDNA prepared from mRNA isolated from a human T-cell leukemic cell line designated CCRF-HSB-2 (ATCC CCL-
30 120.1). The PCR products containing the 5' end of the hek DNA were digested with SpeI and HindIII to isolate a DNA fragment extending from the 5' end of the mature human hek sequence (i.e., lacking DNA encoding the signal sequence) to a HindIII site found in the hek gene. The PCR products containing the 3' end of the hek extracellular domain DNA were digested with HindIII and ClaI to isolate a fragment extending from the
35 internal HindIII site to a ClaI site just downstream of the 3' end of the sequence encoding

the hek extracellular domain. The ClaI site is in a multiple cloning site (mcs) introduced just downstream of the extracellular domain.

DNA encoding a mutein of the Fc region of a human IgG1 antibody was isolated. This Fc mutein DNA and the polypeptide encoded thereby are described in U.S. patent application serial no. 08/097,827, entitled "Novel Cytokine Which is a Ligand for OX40" filed July 23, 1993, which application is incorporated herein by reference. The mutein DNA was derived from a native Fc polypeptide-encoding DNA by site-directed mutagenesis conducted essentially as described by Deng and Nickoloff, *Anal. Biochem.* 200:81 (1992). The amino acid sequence of the Fc mutein polypeptide is identical to that of the native Fc polypeptide described in PCT application WO 93/10151, except that amino acid 19 has been changed from Leu to Ala, amino acid 20 has been changed from Leu to Glu, and amino acid 22 has been changed from Gly to Ala. This mutein Fc exhibits reduced affinity for immunoglobulin receptors.

A recombinant vector containing the Fc mutein DNA was cleaved with ClaI and NotI, which cleave the vector in a polylinker region immediately upstream and downstream, respectively, of the Fc mutein DNA insert. The desired Fc mutein-encoding fragment was isolated.

The mutein Fc polypeptide extends from the N-terminal hinge region to the native C-terminus, i.e., is an essentially full-length antibody Fc region. Fragments of Fc regions, e.g., those that are truncated at the C-terminal end, also may be employed. The fragments preferably contain multiple cysteine residues (at least the cysteine residues in the hinge reaction) to permit interchain disulfide bonds to form between the Fc polypeptide portions of two separate hek:Fc fusion proteins, creating dimers.

A mammalian expression vector designated SMAG4 was cleaved with SpeI and NotI. The SMAG4 vector comprises a murine interleukin-7 signal peptide-encoding sequence (described in U.S. Patent 4,965,195) inserted into the mammalian high expression vector pDC201 (described in Sims et al., *Science* 241:585, 1988, and in PCT application WO 89/03884), which is also capable of replication in *E. coli*. SpeI cleaves the vector immediately downstream of the IL-7 signal peptide-encoding sequence. NotI cleaves approximately 155 bp downstream of the SpeI site in a multiple cloning site of the vector. The large SpeI/NotI fragment containing the vector sequences and the IL-7 signal peptide-encoding DNA was isolated.

A four-way ligation was conducted to insert the two hek-encoding DNA fragments and the Fc mutein-encoding DNA fragment described above into the SpeI/NotI

cleaved SMAG4 expression vector. *E. coli* cells were transfected with the ligation mixture and the desired recombinant vector was isolated therefrom. The isolated vector encodes a fusion protein comprising (from N- to C-terminus) the murine IL-7 signal peptide, the hek extracellular domain, four amino acids encoded by the introduced mcs, and the Fc mutein.

The expression vector was then co-transfected with plasmid pSV3.NEO into CV1/EBNA cells. The CV1/EBNA cell line (ATCC CRL 10478) was derived from a monkey kidney cell line as described in McMahan et al. (*EMBO J.*, 10:2821, 1991). Vector pSV3.NEO expresses SV40 T-antigen, which is not produced by the host cells. The pSV3.NEO vector is similar to pSV3 (Mulligan and Berg, *Proc. Natl. Acad. Sci. USA* 78:2072, 1981), but additionally contains a neomycin resistance gene. The transformed cells were cultivated to allow transient expression of the fusion protein, which is secreted into the culture medium *via* the murine IL-7 signal peptide. The fusion protein can be purified on a protein A Sepharose column, eluted, and used to screen cells for the ability to bind the hek:Fc protein or elk:Fc protein.

EXAMPLE 5: Binding Study

The ability of LERK-6 to bind to elk or hek can be determined by using the following assay. Cells expressing LERK-6 on the cell surface are prepared. LERK-6 DNA is amplified by PCR. The primers employed in the PCR are selected to define the termini of the coding region of the LERK-6 DNA, and also include a Xho I restriction site at the 5' end and a Not I site at the 3' end of the amplified DNA. The 5' primer additionally included a consensus Kozak sequence upstream of the initiation codon.

The reaction products are digested with Xho I and Not I and inserted into an expression vector cleaved with Sal I (which is compatible with Xho I) and Not I. The expression vector can be pDC410, which is a mammalian expression vector that also replicates in *E. coli*. pDC410 is similar to pDC406 (McMahan et al., *EMBO J.* 10:2821, 1991). The pDC410 multiple cloning site (mcs) differs from that of pDC406 in that it contains additional restriction sites and three stop codons (one in each reading frame). A T7 polymerase promoter downstream of the mcs facilitates sequencing of DNA inserted into the mcs. In addition, the EBV origin of replication is replaced by DNA encoding the SV40 large T antigen (driven from an SV40 promoter) in pDC410.

CV1-EBNA-1 cells in 10 cm² dishes are transfected with the recombinant expression vector containing LERK-6 DNA. The CV-1/EBNA-1 cell line (ATCC CRL 10478) constitutively expresses EBV nuclear antigen-1 driven from the CMV immediate-

early enhancer/promoter. CV1-EBNA-1 is derived from the African Green Monkey kidney cell line CV-1 (ATCC CCL 70), as described by McMahan et al. (*EMBO J.* 10:2821, 1991).

5 The transfected cells are cultured for 24 hours, and the cells in each dish then are split into a 24-well plate. After culturing an additional 48 hours, a binding assay is conducted by the following procedure. The transfected cells (about 4×10^4 cells/well) are washed with BM-NFDM, which is binding medium (RPMI 1640 containing 25 mg/ml bovine serum albumin, 2 mg/ml sodium azide, 20 mM Hepes pH 7.2) to which 50 mg/ml nonfat dry milk has been added. The cells then are incubated for 1 hour at 37°C with
10 various concentrations of the elk:Fc fusion protein prepared in Example 2 or the hek:Fc fusion protein prepared in Example 3. Cells then are washed and incubated with a constant saturating concentration of a ^{125}I -mouse anti-human IgG in binding medium, with gentle agitation for 1 hour at 37°C. After extensive washing, cells are released *via* trypsinization.

15 The mouse anti-human IgG employed above is directed against the Fc region of human IgG and can be obtained from Jackson Immunoresearch Laboratories, Inc., West Grove, PA. The antibody is radioiodinated using the standard chloramine-T method. The antibody will bind to the Fc portion of any elk:Fc or hek:Fc fusion protein that has bound to the cells. In all assays, non-specific binding of ^{125}I -antibody is assayed in the absence
20 of elk:Fc (or hek:Fc), as well as in the presence of elk:Fc (or hek:Fc) and a 200-fold molar excess of unlabeled mouse anti-human IgG antibody.

Cell-bound ^{125}I -antibody is quantified on a Packard Autogamma counter. Affinity calculations (Scatchard, *Ann. N.Y. Acad. Sci.* 51:660, 1949) are generated on
25 RS/1 (BBN Software, Boston, MA) run on a Microvax computer.

EXAMPLE 6 **Monoclonal Antibodies to LERK-6**

This example illustrates a method for preparing monoclonal antibodies to LERK-
30 6. Purified LERK-6, a fragment thereof such as the extracellular domain, synthetic peptides or cells that express LERK-6 can be used to generate monoclonal antibodies against LERK-6 using conventional techniques, for example, those techniques described in U.S. Patent 4,411,993. Briefly, mice are immunized with LERK-6 as an immunogen emulsified in complete Freund's adjuvant, and injected in amounts ranging from 10-100
35 µg subcutaneously or intraperitoneally. Ten to twelve days later, the immunized animals are boosted with additional LERK-6 emulsified in incomplete Freund's adjuvant. Mice

are periodically boosted thereafter on a weekly to bi-weekly immunization schedule. Serum samples are periodically taken by retro-orbital bleeding or tail-tip excision to test for LERK-6 antibodies by dot blot assay, ELISA (Enzyme-Linked Immunosorbent Assay) or inhibition of hek or elk binding.

5

Following detection of an appropriate antibody titer, positive animals are provided one last intravenous injection of LERK-6 in saline. Three to four days later, the animals are sacrificed, spleen cells harvested, and spleen cells are fused to a murine myeloma cell line, e.g., NS1 or preferably P3x63Ag8.653 (ATCC CRL 1580). Fusions generate
10 hybridoma cells, which are plated in multiple microtiter plates in a HAT (hypoxanthine, aminopterin and thymidine) selective medium to inhibit proliferation of non-fused cells, myeloma hybrids, and spleen cell hybrids.

The hybridoma cells are screened by ELISA for reactivity against purified LERK-
15 6 by adaptations of the techniques disclosed in Engvall et al., *Immunochem.* 8:871, 1971 and in U.S. Patent 4,703,004. A preferred screening technique is the antibody capture technique described in Beckmann et al., (*J. Immunol.* 144:4212, 1990) Positive hybridoma cells can be injected intraperitoneally into syngeneic BALB/c mice to produce ascites containing high concentrations of anti-LERK-6-L monoclonal antibodies.
20 Alternatively, hybridoma cells can be grown *in vitro* in flasks or roller bottles by various techniques. Monoclonal antibodies produced in mouse ascites can be purified by ammonium sulfate precipitation, followed by gel exclusion chromatography. Alternatively, affinity chromatography based upon binding of antibody to protein A or protein G can also be used, as can affinity chromatography based upon binding to LERK-
25 6.

EXAMPLE 7

Generation of Transgenic Mice That Overexpress LERK-6

This example describes a procedure for generating transgenic mice that
30 overexpress LERK-6. LERK-6-overexpressing transgenic mice can be studied to determine the biological effects of overexpression. Mouse (B16/J) pronuclei are microinjected with LERK-6 DNA according to the method described by Gordon et al., *Science* 214:1244-1246, (1981). In general, fertilized mouse eggs having visible pronuclei are first placed on an injection chamber and held in place with a small pipet.
35 An injection pipet is used to inject the gene encoding the LERK-6 (e.g., clone #6C) into the pronuclei of the egg. Injected eggs are either (i) transferred into the oviduct of a 0.5 day p.c. pseudopregnant female; (ii) cultured *in vitro* to the two-cell stage (overnight) and

transferred into the oviduct of a 0.5 day p.c. pseudopregnant female; or (iii) cultured *in vitro* to the blastocyst stage and transferred into the uterus of a 2.5 day p.c. pseudopregnant female. Preferably, either of the first two options can be used since they avoid extended *in vitro* culture, and preferably, approximately 20-30 microinjected eggs
5 should be transferred to avoid small litters.

What is claimed is:

1. An isolated DNA sequence encoding a LERK-6 polypeptide that binds hek/elk.
2. An isolated DNA sequence encoding a LERK-6 polypeptide that binds hek/elk and that is at least 90% identical with the sequence of amino acid residues selected from the group consisting of 1 to 184 of SEQ ID NO:2, 1 to 104 of SEQ ID NO:8 and 1 to 184 of SEQ ID NO:10.
3. An isolated DNA sequence according to claim 1, comprising the nucleotide sequence selected from the group consisting of 1 to 552 of SEQ ID NO:1, 2 to 313 of SEQ ID NO:7 and 88 to 642 of SEQ ID NO:9.
4. A DNA according to claim 1, selected from the group consisting of:
 - (a) cDNA sequences selected from the group consisting of SEQ ID NO:1, SEQ ID NO:7, and nucleotides 88 to 642 of SEQ ID NO:9 that code for LERK-6;
 - (b) DNA sequences that hybridize under highly stringent conditions to the cDNA of (a); and which DNA sequences encode LERK-6; and
 - (c) DNA sequences that, due to the degeneracy of the genetic code, encode LERK-6 polypeptides having the amino acid sequence of the polypeptides encoded by the DNA sequences of (a) or (b).
5. An isolated and purified LERK-6 polypeptide that binds hek/elk.
6. An isolated and purified LERK-6 polypeptide that comprises an amino acid sequence that is at least 90% identical to the sequence of amino acid residues selected from the group consisting of 1 to 184 of SEQ ID NO:2, 1 to 104 of SEQ ID NO:8 and 1 to 184 of SEQ ID NO:10.
7. A LERK-6 polypeptide according to claim 5, encoded by the nucleotide sequence according to claim 4.
8. An isolated and purified LERK-6 polypeptide that comprises an amino acid sequence that is at least 90% identical to the sequence of amino acid residues selected from the group consisting of 1 to 145 of SEQ ID NO:2, 1 to 104 of SEQ ID NO:8, and 1 to 145 of SEQ ID NO:10.

9. An isolated and purified LERK-6 polypeptide that comprises the amino acid residues 1 to 145 of SEQ ID NO:2.
10. An isolated and purified LERK-6 polypeptide comprising the amino acid residues 1 to 184 of SEQ ID NO:10.
11. A polypeptide according to claim 5 and selected from the group consisting of polypeptides encoded by the cDNA insert of vector λ 13MLERK-6 λ gt10 having accession number ATCC 75829 and polypeptides encoded by the cDNA insert of the vector having accession number_____.
12. An expression vector comprising a DNA sequence according to claim 1.
13. An expression vector comprising a DNA sequence according to claim 2.
14. An expression vector comprising a DNA sequence according to claim 3.
15. An expression vector comprising a DNA sequence according to claim 4.
16. A host cell transfected or transformed with the expression vector according to claim 12.
17. A host cell transformed or transfected with the expression vector according to claim 13.
18. A host cell transformed or transfected with the expression vector according to claim 14.
19. A host cell transformed or transfected with the expression vector according to claim 14.
20. A process for producing a LERK-6 polypeptide, comprising culturing a host cell according to claim 16 under conditions promoting expression, and recovering the polypeptide from the culture medium.

21. A process for producing a LERK-6 polypeptide, comprising culturing a host cell according to claim 17 under conditions promoting expression, and recovering the polypeptide from the culture medium.
22. A process for producing a LERK-6 polypeptide, comprising culturing a host cell according to claim 18 under conditions promoting expression, and recovering the polypeptide from the culture medium.
23. A process for producing a LERK-6 polypeptide, comprising culturing a host cell according to claim 19 under conditions promoting expression, and recovering the polypeptide from the culture medium.
24. An isolated and purified antibody that is immunoreactive with a LERK-6 polypeptide.
25. An antibody according to claim 24 that is a monoclonal antibody.
26. A transgenic non-human mammal all of whose germ and somatic cells contain a DNA sequence according to claim 1 introduced into said mammal, or an ancestor of said mammal, at an embryonic stage.
27. A method of separating cells having the hek/elk receptor on the surface thereof from a mixture of cells in suspension, comprising contacting the cells in the mixture with a contacting surface having a LERK-6 polypeptide according to claim 5 thereon, and separating the contacting surface and the suspension.
28. A method for delivering a desired molecule to a cell having hek/elk on its surface, comprising contacting the hek/elk with a fusion protein comprising a LERK-6 polypeptide according to claim 5 and the desired molecule.

1

- (i) APPLICANT: IMMUNEX CORPORATION
- (ii) TITLE OF INVENTION: Cytokine Designated as LERK-6
- (iii) NUMBER OF SEQUENCES: 8
- (iv) CORRESPONDENCE ADDRESS:
 - (A) ADDRESSEE: Immunex Corporation
 - (B) STREET: 51 University Street
 - (C) CITY: Seattle
 - (D) STATE: WA
 - (E) COUNTRY: USA
 - (F) ZIP: 98101
- (v) COMPUTER READABLE FORM:
 - (A) MEDIUM TYPE: Floppy disk
 - (B) COMPUTER: IBM/PC Compatible
 - (C) OPERATING SYSTEM: MS-DOS/Windows 95
 - (D) SOFTWARE: Word for Windows, Version 7.0a
- (vi) CURRENT APPLICATION DATA:
 - (A) APPLICATION NUMBER: PCT --to be assigned--
 - (B) FILING DATE: 27 August 1998
 - (C) CLASSIFICATION:
- (viii) ATTORNEY/AGENT INFORMATION:
 - (A) NAME: Henry, Janis C.
 - (B) REGISTRATION NUMBER: 34,347
 - (C) REFERENCE/DOCKET NUMBER: 2826
- (ix) TELECOMMUNICATION INFORMATION:
 - (A) TELEPHONE: (206) 587-0430
 - (B) TELEFAX: (206) 233-0644

```

(i) SEQUENCE CHARACTERISTICS:
    (A) LENGTH: 555 base pairs
    (B) TYPE: nucleic acid
    (C) STRANDEDNESS: single
    (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(vii) IMMEDIATE SOURCE:
    (B) CLONE: LERK-6

(ix) FEATURE:
    (A) NAME/KEY: CDS
    (B) LOCATION: 1..552

```

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:

GCC CGG GCC AAC GCT GAC CGA TAC GCA GTC TAC TGG AAC CGT AGC AAC
Ala Arg Ala Asn Ala Asp Arg Tyr Ala Val Tyr Trp Asn Arg Ser Asn
1 5 10 15

CCC Pro	AGG Arg	TTT Phe	CAG Gln 20	GTG Val	AGC Ser	GCT Ala	GTG Val	GGT Gly 25	GAT Asp	GGC Gly	GGC Gly	GGC Gly	TAT Tyr 30	ACC Thr	GTG Val	96
GAG Glu	GTG Val	AGC Ser 35	ATC Ile	AAC Asn	GAC Asp	TAC Tyr	CTG Leu 40	GAT Asp	ATC Ile	TAC Tyr	TGC Cys	CCA Pro 45	CAC His	TAC Tyr	GGG Gly	144
GCG Ala	CCG Pro 50	CTG Leu	CCC Pro	CCG Pro	GCT Ala	GAG Glu 55	CGC Arg	ATG Met	GAG Glu	CGG Arg	TAC Tyr 60	ATC Ile	CTG Leu	TAC Tyr	ATG Met	192
GTG Val 65	AAT Asn	GGT Gly	GAG Glu	GGC Gly	CAC His 70	GCC Ala	TCC Ser	TGT Cys	GAC Asp	CAC His 75	CGG Arg	CAG Gln	CGA Arg	GGC Gly	TTC Phe 80	240
AAG Lys	CGC Arg	TGG Trp	GAA Glu	TGC Cys 85	AAC Asn	CGG Arg	CCC Pro	GCA Ala	GCG Ala 90	CCC Pro	GGG Gly	GGA Gly	CCC Pro	CTC Leu 95	AAG Lys	288
TTC Phe	TCA Ser	GAG Glu	AAG Lys 100	TTC Phe	CAA Gln	CTC Leu	TTC Phe	ACC Thr 105	CCC Pro	TTT Phe	TCC Ser	CTG Leu 110	GGC Gly	TTT Phe	GAG Glu	336
TTC Phe	CGG Arg	CCT Pro 115	GGC Gly	CAC His	GAA Glu	TAC Tyr	TAC Tyr 120	TAC Tyr	ATC Ile	TCT Ser	GCC Ala	ACA Thr 125	CCT Pro	CCC Pro	AAC Asn	384
CTC Leu	GTG Val 130	GAC Asp	CGA Arg	CCC Pro	TGC Cys	CTG Leu 135	CGA Arg	CTC Leu	AAG Lys	GTT Val	TAT Tyr 140	GTG Val	CGT Arg	CCA Pro	ACC Thr	432
AAT Asn 145	GAG Glu	ACC Thr	CTG Leu	TAT Tyr	GAG Glu 150	GCT Ala	CCA Pro	GAG Glu	CCC Pro 155	ATC Ile	TTC Phe	ACC Thr	AGT Ser	AAC Asn	AGC Ser 160	480
TCC Ser	TGC Cys	AGC Ser	GGC Gly	CTG Leu 165	GGT Gly	GGC Gly	TGC Cys	CAC His	CTC Leu 170	TTC Phe	CTC Leu	ACC Thr	ACC Thr	GTC Val 175	CCT Pro	528
GTG Val	CTG Leu	TGG Trp	TCC Ser 180	CTT Leu	CTG Leu	GGC Gly	TCC Ser	TAG								555

(2) INFORMATION FOR SEQ ID NO:2:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 184 amino acids
(B) TYPE: amino acid
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: protein

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:

Ala	Arg	Ala	Asn	Ala	Asp	Arg	Tyr	Ala	Val	Tyr	Trp	Asn	Arg	Ser	Asn
1				5					10					15	
Pro	Arg	Phe	Gln	Val	Ser	Ala	Val	Gly	Asp	Gly	Gly	Gly	Tyr	Thr	Val
			20					25					30		
Glu	Val	Ser	Ile	Asn	Asp	Tyr	Leu	Asp	Ile	Tyr	Cys	Pro	His	Tyr	Gly
		35					40					45			
Ala	Pro	Leu	Pro	Pro	Ala	Glu	Arg	Met	Glu	Arg	Tyr	Ile	Leu	Tyr	Met
	50					55					60				

Val Asn Gly Glu Gly His Ala Ser Cys Asp His Arg Gln Arg Gly Phe
65 70 75 80

Lys Arg Trp Glu Cys Asn Arg Pro Ala Ala Pro Gly Gly Pro Leu Lys
85 90 95

Phe Ser Glu Lys Phe Gln Leu Phe Thr Pro Phe Ser Leu Gly Phe Glu
100 105 110

Phe Arg Pro Gly His Glu Tyr Tyr Tyr Ile Ser Ala Thr Pro Pro Asn
115 120 125

Leu Val Asp Arg Pro Cys Leu Arg Leu Lys Val Tyr Val Arg Pro Thr
130 135 140

Asn Glu Thr Leu Tyr Glu Ala Pro Glu Pro Ile Phe Thr Ser Asn Ser
145 150 155 160

Ser Cys Ser Gly Leu Gly Gly Cys His Leu Phe Leu Thr Thr Val Pro
165 170 175

Val Leu Trp Ser Leu Leu Gly Ser
180

(2) INFORMATION FOR SEQ ID NO:3:

- (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 28 base pairs
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:

GATATTTACT GCCCGCACTA CAACAGCT

28

(2) INFORMATION FOR SEQ ID NO:4:

- (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 25 base pairs
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:

AGAGAAGGCG CTGTAGCGCT GGAAC

25

(2) INFORMATION FOR SEQ ID NO:5:

- (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 31 base pairs
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

- (ii) MOLECULE TYPE: cDNA
- (iii) HYPOTHETICAL: NO
- (iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:5:

ACGTAGTCTA CTGGAAGTCC AGTAACCCCA G

31

(2) INFORMATION FOR SEQ ID NO:6:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 34 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: cDNA
- (iii) HYPOTHETICAL: NO
- (iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:6:

AGCCTCAAGC ACTGGCCAGA ACTCTCTCTG GAGT

34

(2) INFORMATION FOR SEQ ID NO:7:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 314 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: cDNA to mRNA
- (iii) HYPOTHETICAL: NO
- (iv) ANTI-SENSE: NO

(ix) FEATURE:

- (A) NAME/KEY: CDS
- (B) LOCATION: 2..313

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:7:

G	TTC	CAC	GCA	GGC	GCG	GGG	GAC	GAC	GGC	GGG	GGC	TAC	ACG	GTG	GAG	46
	Phe	His	Ala	Gly	Ala	Gly	Asp	Asp	Gly	Gly	Gly	Tyr	Thr	Val	Glu	
1				5						10					15	
GTG	AGC	ATC	AAT	GAC	TAC	CTG	GAC	ATC	TAC	TGC	CCG	CAC	TAT	GGG	GCG	94
Val	Ser	Ile	Asn	Asp	Tyr	Leu	Asp	Ile	Tyr	Cys	Pro	His	Tyr	Gly	Ala	
				20					25					30		
CCG	CTG	CCG	CCG	GCC	GAG	CGC	ATG	GAG	CAC	TAC	GTG	CTG	TAC	ATG	GTC	142
Pro	Leu	Pro	Pro	Ala	Glu	Arg	Met	Glu	His	Tyr	Val	Leu	Tyr	Met	Val	
				35				40						45		

5

```

AAC GGC GAG GGC CAC GCC TCC TGC GAC CAC CGC CAG CGC GGC TTC AAG 190
Asn Gly Glu Gly His Ala Ser Cys Asp His Arg Gln Arg Gly Phe Lys
      50                      55                      60

CGC TGG GAG TGC AAC CGG CCC GCG GCG CCC GGG GGG CCG CTC AAG TTC 238
Arg Trp Glu Cys Asn Arg Pro Ala Ala Pro Gly Gly Pro Leu Lys Phe
      65                      70                      75

TCG GAG AAG TTC CAG CTC TTC ACG CCC TTC TCC CTG GGC TTC GAG TTC 286
Ser Glu Lys Phe Gln Leu Phe Thr Pro Phe Ser Leu Gly Phe Glu Phe
      80                      85                      90

CGG CCC GGC CAC GAG TAT TAC TAC ATC T 314
Arg Pro Gly His Glu Tyr Tyr Tyr Ile
      100

```

(2) INFORMATION FOR SEQ ID NO:8:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 104 amino acids
- (B) TYPE: amino acid
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: protein

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:8:

```

Phe His Ala Gly Ala Gly Asp Asp Gly Gly Gly Tyr Thr Val Glu Val
 1              5              10              15

Ser Ile Asn Asp Tyr Leu Asp Ile Tyr Cys Pro His Tyr Gly Ala Pro
      20              25              30

Leu Pro Pro Ala Glu Arg Met Glu His Tyr Val Leu Tyr Met Val Asn
      35              40              45

Gly Glu Gly His Ala Ser Cys Asp His Arg Gln Arg Gly Phe Lys Arg
      50              55              60

Trp Glu Cys Asn Arg Pro Ala Ala Pro Gly Gly Pro Leu Lys Phe Ser
      65              70              75              80

Glu Lys Phe Gln Leu Phe Thr Pro Phe Ser Leu Gly Phe Glu Phe Arg
      85              90              95

Pro Gly His Glu Tyr Tyr Tyr Ile
      100

```

(2) INFORMATION FOR SEQ ID NO:9:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 642 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA to mRNA

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(ix) FEATURE:

- (A) NAME/KEY: CDS
- (B) LOCATION: 1...642

(A) NAME/KEY: Signal Peptide
(B) LOCATION: 1-90

ATG GCG CCC GCG CAG CGC CCG CTG CTC CCG CTG CTG CTC CTG CTG TTA	48
Met Ala Pro Ala Gln Arg Pro Leu Leu Pro Leu Leu Leu Leu Leu	
-25 -20 -15	
CCG CTG CCG CCG CCG CCC TTC GCG CGC GCC GAG GAC GCC GCC CGC GCC	96
Pro Leu Pro Pro Pro Phe Ala Arg Ala Glu Asp Ala Ala Arg Ala	
-10 -5 1	
AAC TCG GAC CGC TAC GCC GTC TAC TGG AAC CGC AGC AAC CCC AGG TTC	144
Asn Ser Asp Arg Tyr Ala Val Tyr Trp Asn Arg Ser Asn Pro Arg Phe	
5 10 15	
CAC GCA GGC GCG GGG GAC GAC GGC GGG GGC TAC ACG GTG GAG GTG AGC	192
His Ala Gly Ala Gly Asp Asp Gly Gly Gly Tyr Thr Val Glu Val Ser	
20 25 30 35	
ATC AAT GAC TAC CTG GAC ATC TAC TGC CCG CAC TAT GGG GCG CCG CTG	240
Ile Asn Asp Tyr Leu Asp Ile Tyr Cys Pro His Tyr Gly Ala Pro Leu	
40 45 50	
CCG CCG GCC GAG CGC ATG GAG CAC TAC GTG CTG TAC ATG GTC AAC GGC	288
Pro Pro Ala Glu Arg Met Glu His Tyr Val Leu Tyr Met Val Asn Gly	
55 60 65	
GAG GGC CAC GCC TCC TGC GAC CAC CGC CAG CGC GGC TTC AAG CGC TGG	336
Glu Gly His Ala Ser Cys Asp His Arg Gln Arg Gly Phe Lys Arg Trp	
70 75 80	
GAG TGC AAC CGG CCC GCG GCG CCC GGG GGG CCG CTC AAG TTC TCG GAG	384
Glu Cys Asn Arg Pro Ala Ala Pro Gly Gly Pro Leu Lys Phe Ser Glu	
85 90 95	
AAG TTC CAG CTC TTC ACG CCC TTC TCC CTG GGC TTC GAG TTC CGG CCC	432
Lys Phe Gln Leu Phe Thr Pro Phe Ser Leu Gly Phe Glu Phe Arg Pro	
100 105 110 115	
GGC CAC GAG TAT TAC TAC ATC TCT GCC ACG CCT CCC AAT GCT GTG GAC	480
Gly His Glu Tyr Tyr Tyr Ile Ser Ala Thr Pro Pro Asn Ala Val Asp	
120 125 130	
CGG CCC TGC CTG CGA CTG AAG GTG TAC GTG CGG CCG ACC AAC GAG ACC	528
Arg Pro Cys Leu Arg Leu Lys Val Tyr Val Arg Pro Thr Asn Glu Thr	
135 140 145	
CTG TAC GAG GCT CCT GAG CCC ATC TTC ACC AGC AAT AAC TCG TGT AGC	576
Leu Tyr Glu Ala Pro Glu Pro Ile Phe Thr Ser Asn Asn Ser Cys Ser	
150 155 160	
AGC CCG GGC GGC TGC CGC CTC TTC CTC AGC ACC ATC CCC GTG CTC TGG	624
Ser Pro Gly Gly Cys Arg Leu Phe Leu Ser Thr Ile Pro Val Leu Trp	
165 170 175	
ACC CTC CTG GGT TCC TAG	642
Thr Leu Leu Gly Ser *	
180	

(2) INFORMATION FOR SEQ ID NO:10:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 214 amino acids

(B) TYPE: amino acid

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: protein

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:10:

[illegible]

INTERNATIONAL SEARCH REPORT

National Application No.

PCT/US 98/17772

A. CLASSIFICATION OF SUBJECT MATTER

IPC 6 C12N15/19 C07K14/52 C07K16/24 C07K19/00 A01K67/027

According to International Patent Classification (IPC) or to both national classification and IPC

B. FIELDS SEARCHED

Minimum documentation searched (classification system followed by classification symbols)

IPC 6 C07K C12N

Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched

Electronic data base consulted during the international search (name of data base and, where practical, search terms used)

C. DOCUMENTS CONSIDERED TO BE RELEVANT

Category *	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
X	WO 96 10911 A (IMMUNEX CORP) 18 April 1996 see the whole document ----	1-28
A	WO 96 17925 A (IMMUNEX CORP) 13 June 1996 see figures SEQ.ID.4,5; example 1 ----	1-28
A	WO 96 09384 A (HARVARD COLLEGE) 28 March 1996 see abstract; claim 1 ----	1-28
P,X	AASHEIM HC ET AL: "Cloning, chromosomal mapping, and tissue expression of the gene encoding the human Eph-family kinase ligand ephrin-A2." BIOCHEM BIOPHYS RES COMMUN, NOV 18 1998, 252 (2) P378-82, XP002090332 UNITED STATES see the whole document -----	1-25

☐ Further documents are listed in the continuation of box C.

☒ Patent family members are listed in annex.

* Special categories of cited documents:

- "A" document defining the general state of the art which is not considered to be of particular relevance
- "E" earlier document but published on or after the international filing date
- "L" document which may throw doubts on priority claim(s) or which is cited to establish the publication date of another citation or other special reason (as specified)
- "O" document referring to an oral disclosure, use, exhibition or other means
- "P" document published prior to the international filing date but later than the priority date claimed

- "T" later document published after the international filing date or priority date and not in conflict with the application but cited to understand the principle or theory underlying the invention
- "X" document of particular relevance; the claimed invention cannot be considered novel or cannot be considered to involve an inventive step when the document is taken alone
- "Y" document of particular relevance; the claimed invention cannot be considered to involve an inventive step when the document is combined with one or more other such documents, such combination being obvious to a person skilled in the art.
- "&" document member of the same patent family

Date of the actual completion of the international search

20 January 1999

Date of mailing of the international search report

01/02/1999

Name and mailing address of the ISA

European Patent Office, P.B. 5818 Patentlaan 2
NL - 2280 HV Rijswijk
Tel. (+31-70) 340-2040, Tx. 31 651 epo nl,
Fax: (+31-70) 340-3016

Authorized officer

Gurdjian, D

INTERNATIONAL SEARCH REPORT

International application No.:

PCT/US 93/17772

Box I Observations where certain claims were found unsearchable (Continuation of item 1 of first sheet)

This International Search Report has not been established in respect of certain claims under Article 17(2)(a) for the following reasons:

1. ☒ Claims Nos.:
because they relate to subject matter not required to be searched by this Authority, namely:
Remark: Although claim 28 as far as it relates to an in vivo method is directed to a method of treatment of the human/animal body, the search has been carried out and based on the alleged effects of the compound/
composition.
2. ☐ Claims Nos.:
because they relate to parts of the International Application that do not comply with the prescribed requirements to such an extent that no meaningful International Search can be carried out, specifically:
3. ☐ Claims Nos.:
because they are dependent claims and are not drafted in accordance with the second and third sentences of Rule 6.4(a).

Box II Observations where unity of invention is lacking (Continuation of item 2 of first sheet)

This International Searching Authority found multiple inventions in this international application, as follows:

1. ☐ As all required additional search fees were timely paid by the applicant, this International Search Report covers all searchable claims.
2. ☐ As all searchable claims could be searched without effort justifying an additional fee, this Authority did not invite payment of any additional fee.
3. ☐ As only some of the required additional search fees were timely paid by the applicant, this International Search Report covers only those claims for which fees were paid, specifically claims Nos.:
4. ☐ No required additional search fees were timely paid by the applicant. Consequently, this International Search Report is restricted to the invention first mentioned in the claims; it is covered by claims Nos.:

Remark on Protest

☐ The additional search fees were accompanied by the applicant's protest.

☐ No protest accompanied the payment of additional search fees.

INTERNATIONAL SEARCH REPORT

Information on patent family members

International Application No

PCT/US 98/17772

Patent document cited in search report	Publication date	Patent family member(s)	Publication date
WO 9610911 A	18-04-1996	AU 687647 B	26-02-1998
		AU 3826895 A	02-05-1996
		EP 0785716 A	30-07-1997
		FI 971276 A	27-05-1997
		NO 971378 A	05-06-1997
WO 9617925 A	13-06-1996	AU 4639396 A	26-06-1996
		CA 2206488 A	13-06-1996
		EP 0871702 A	21-10-1998
		FI 972390 A	05-06-1997
		JP 10512440 T	02-12-1998
		NO 972455 A	06-08-1997
WO 9609384 A	28-03-1996	AU 3592095 A	09-04-1996
		US 5795734 A	18-08-1998